

Helbing Laboratory eDNA Technical Bulletin

All eDNA tools are validated through a rigorous multi-step evaluation protocol that includes tests of DNA target specificity and amplification sensitivity¹⁻³.

| General eDNA Assay Information | | | | | | |
|--|--------------------------|-----------------------|--|--|--|--|
| Target Species: Chloroplasts (Plant/Algae) | eDNA qPCR Tool: ePlant5 | Gene Target: 23S | | | | |
| Species Code: IntegritE-DNA | eDNA qPCR Format: TaqMan | Published in: 1, 2, 5 | | | | |

| eDNA Assay Sensitivity Test Summary using gBlocks™ Synthetic DNA | | | | | | | | | | | | | |
|--|-----|-----|--|-----|--------|------------------|-----|--------|-----|--------|-----|-----|--------|
| | LOD | N/A | 95% CI | N/A | Copies | LOQ | N/A | 95% CI | N/A | Copies | LOB | N/A | hits/8 |
| Binomial-Poisson model for 8 technical replicates | | | | | | | | | | | | | |
| Determined using eLowQuant R code ⁴ . | | | When the LOQ < LOD, use the LOD for the LOQ. | | | Enzyme: Immolase | | | | | | | |

eDNA Assay Specificity Test Information

Each qPCR reaction in the specificity assay contained 10 picograms of voucher target gDNA (n=25 technical replicates)

Chloroplast DNA is used as an endogenous quality control for eDNA samples when testing for animal DNA.

A broad range of plants and algae from fresh water, marine water, sediments, air, and soil have been tested and pass.

Fresh, unopened bottled water typically does not have plant or algae DNA in it.

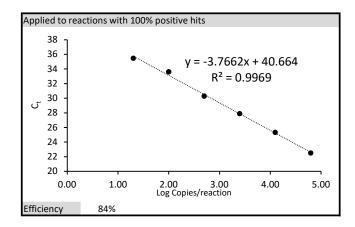
Typically the C_{α} cutoff is 27 for pass/fail for most applications.

"Fail" indicates poor quality sample with inhibition and/or degradation.

Refer fo references 1 and 2 for application examples.

The continuous standard curve is based upon n=8 technical replicates with very small standard error.

Based on a 2 μL DNA input in a total 15 μL reaction



References

- 1. Hobbs, J, Adams, IT, Round, JM, Goldberg, CS, Allison, MJ, Bergman, LC, Mirabzadeh, A, Allen, H, Helbing, CC (2020) Revising the range of Rocky Mountain tailed frog, Ascaphus montanus, in British Columbia, Canada, using environmental DNA methods. Environmental DNA, 2: 350-361. https://doi.org/10.1002/edn3.82
- 2. Hobbs, J, Round, JM, Allison, MJ, Helbing, CC (2019) Expansion of the known distribution of the coastal tailed frog, Ascaphus truei, in British Columbia, Canada, using robust eDNA detection methods. PLOS ONE 14(3): e0213849. https://doi.org/10.1371/journal.pone.0213849
- 3. Langlois, VS, Allison, MJ, Bergman, LC, To, TA, and Helbing, CC (2020) The need for robust qPCR-based eDNA detection assays in environmental monitoring and risk assessments. Environmental DNA, 3: 519-527. doi: 10.1002/edn3.164
- 4. Lesperance, M, Allison, MJ, Bergman, LC, Hocking, MD, and Helbing, CC (2021) A statistical model for calibration and computation of detection and quantification limits for low copy number environmental DNA samples. Environmental DNA, 3: 970-981. doi: 10.1002/edn3.220
- 5. Veldhoen, N., Hobbs, J., Ikonomou, G., Hii, M., Lesperance, M., and Helbing, C.C. 2016. Implementation of novel design features for qPCR-based eDNA assessment. PLoS One, 11: e0164907. DOI: 10.1371/journal.pone.0164907

| | | Abbreviations | |
|------|---|---------------|--|
| 23S | 23S ribosomal RNA | NTC | qPCR no template control |
| eDNA | Environmental DNA | qPCR | Quantitative real-time polymerase chain reaction |
| gDNA | Total genomic DNA extracted from voucher specimen | SE | Standard error |